

## 特許協力条約に基づいて公開された国際出願

(51) 国際特許分類 6

C07K 14/52, C12N 15/00, 1/21,  
1/19, 5/00, C12P 21/02, C07K 16/24

A1

(11) 国際公開番号

WO 95/13293

(43) 国際公開口

1995年5月18日 (18.05.95)

(21) 国際出願番号

PCT/JP94. 01899

(22) 国際出願日

1994年11月10日 (10. 11. 94)

(30) 優先権データ

特願平5. 305975	1993年11月10日 (10. 11. 93)	JP
特願平5. 342526	1993年12月13日 (13. 12. 93)	JP
特願平6. 74344	1994年3月18日 (18. 03. 94)	JP
特願平6. 180955	1994年7月8日 (08. 07. 94)	JP
特願平6. 239363	1994年9月7日 (07. 09. 94)	JP
特願平6. 278378	1994年10月18日 (18. 10. 94)	JP

中村 純夫 (NAKAMURA, Norio)

〒160 東京都新宿区四谷1丁目7番地

持田製薬株式会社内 Tokyo, (JP)

(74) 代理人

弁理士 渡辺 望松, 外 (WATANABE, Mochitoshi et al.)

〒101 東京都千代田区岩本町3丁目2番2号 千代田岩本ビル4階  
Tokyo, (JP)

(81) 指定国

AU, CA, CN, JP, KR.

(71) 出願人

持田製薬株式会社

(MOCHIDA PHARMACEUTICAL CO., LTD.) (JP/JP)

〒160 東京都新宿区四谷1丁目7番地 Tokyo, (JP)

財団法人 大阪バイオサイエンス研究所

(OSAKA BIOSCIENCE INSTITUTE) (JP/JP)

〒565 大阪府吹田市古江台6丁目2番4号 Osaka, (JP)

(72) 発明者

長田 重一 (NAGATA, Shigekazu)

〒565 大阪府吹田市佐井寺2丁目21-17-511 Osaka, (JP)

須田 貴司 (SUDA, Takashi)

〒562 大阪府箕面市小野原東3丁目4-20-302 Osaka, (JP)

高橋 智浩 (TAKAHASHI, Tomohiro)

添付公開書類

国際調査報告書

**AML** INFORMATION SERVICES  
P.O. BOX 405, CORTE MADERA, CA 94976-0405  
(415) 927-0340 • FAX (415) 927-7250

(54) Title : Fas LIGAND, PART THEREOF, AND DNA CODING FOR THE SAME

(54) 発明の名称 Fasリガンドおよびその一部、およびそれをコードするDNA

(57) Abstract

A Fas ligand useful in the medicinal field and a part thereof; a novel DNA coding for the same; an antibody; a recombinant DNA molecule containing the above DNA; a transformant; a method of purifying the novel protein; a process for producing the novel protein; a novel protein antibody; an oligonucleotide complementary to a Fas ligand gene; and a method of screening a substance associated with the Fas ligand.

a

PMA+イオノマイシン - + + - +

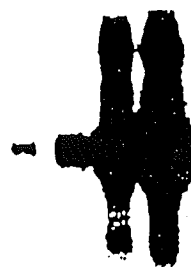
28S ▶

18S ▶

28S ▶

18S ▶

d10S  
d10S  
d10S-2  
d10S-16  
d10S-16



◀ Fas-L



◀ EF1α

a: PMA plus ionomycin

P. 205

Scope of Claims: Claim Articles

1. New polypeptides characterized by the amino acid sequence in the following formula 1 (arrangement number 1 of the sequence lists)

Formula 1

P. 206

2. New polypeptides characterized by the amino acid sequence in the following formula 2 (arrangement number 2 of the sequence lists))

Formula 2

P. 207

3. New polypeptides characterized by the amino acid sequence in the following formula 3 (arrangement number 3 of the sequence lists)

Formula 3

P. 209

4. New polypeptides characterized by the amino acid sequence in the following formula 4 (arrangement number 4 of the sequence lists)

Formula 4

P. 212

5. New polypeptides characterized by the amino acid sequence in the following formula 5 (arrangement number 5 of the sequence lists))

Formula 5

P. 213

6. New polypeptides characterized by the amino acid sequence in the following formula 6 (arrangement number 6 of the sequence lists))

Formula 6

P.215

7. New polypeptides characterized by the amino acid sequence in the following formula 7 (arrangement number 7 of the sequence lists)

Formula 7

P. 217

8. New polypeptides characterized by the amino acid sequence in the following formula 8 (arrangement number 8 of the sequence lists)

Formula 8

P. 219

9. New polypeptides characterized by the amino acid sequence in the following formula 9 (arrangement number 9 of the sequence lists))

Formula 9

P. 221

10. New polypeptides characterized by the amino acid sequence in the following formula 10 (arrangement number 10 of the sequence lists)  
Formula 10

P. 222

11. New polypeptides characterized by the amino acid sequence in the following formula 11 (arrangement number 11 of the sequence lists)  
Formula 11

P. 224

12. New polypeptides characterized by the amino acid sequence in the following formula 12 (arrangement number 12 of the amino acid sequences)  
Formula 12

P. 226-7

13. New polypeptides characterized by the following; contains the amino acid arrangements that are coded by the base sequences that hybridize with the complementary chains of the base arrangements that code any or some of the amino acid sequences chosen from prior formulas 1, 5, and 9. (arrangement numbers 1, 5, and 9 of the sequence lists)

P. 227

14. New polypeptides with the following characteristic: contains the amino acid arrangements coded by the base sequences that hybridize with the complementary chains of the base sequences chosen from any or some of the arrangement numbers 13, 17, and 21 of the sequence lists.

15. Polypeptides recorded in either of claim articles 1 or 14 which bind with Fas antigen.

16. Polypeptides recorded in either claim articles 1 or 15 which have the activity that derives apoptosis towards cells which express Fas antigen.

17. New polypeptides recorded in either claim articles 1 or 16 that are rearranged polypeptides.

18. Polypeptides composed from a part of the amino acid sequences recorded in prior formula 4 (arrangement number 4 of the sequence lists)

19. Polypeptides composed from a part of the amino acid sequences recorded in prior formula 8 (arrangement number 8 of the sequence lists).

20. Polypeptides composed from a part of the amino acid sequences recorded in prior formula 12 (arrangement number 12 of the sequence lists).

21. Polypeptides produced by combining more than 2 kinds of polypeptides that are made from either all or a part of the amino acid sequences chosen from any or some of prior formulas 4, 8, and 12 (arrangement numbers 4, 8, and 12 of the sequence lists).

P. 227-8

22. Polypeptides recorded in either of claim articles 18 or 21 which bind with Fas antigen.

P. 228

23. New DNA which has the characteristic of containing the base sequences that code the new polypeptides recorded in either of claim articles 1 or 22.

24. New DNA which is recorded in claim article 23 in which the aforementioned DNA contains the base sequences which are recorded in the following formula 13 (arrangement number 13 of the sequence lists.)  
Formula 13

P. 229

25. New DNA which is recorded in claim article 23 in which the aforementioned DNA contains the base sequences which are recorded in the following formula 14 (arrangement number 14 of the sequence lists.)  
Formula 14

P. 230-1

26. New DNA which is recorded in claim article 23 in which the aforementioned DNA contains the base sequences which are recorded in the following formula 15 (arrangement number 15 of the sequence lists.)  
Formula 15

P. 232

27. New DNA which is recorded in claim article 23 in which the aforementioned DNA contains the base sequences which are recorded in the following formula 16 (arrangement number 16 of the sequence lists.)  
Formula 16

P. 235

28. New DNA which is recorded in claim article 23 in which the aforementioned DNA contains the base sequences which are recorded in the following formula 17 (arrangement number 17 of the sequence lists.)  
Formula 17

P. 236-7

29. New DNA which is recorded in claim article 23 in which the aforementioned DNA contains the base sequences which are recorded in the following formula 18

(arrangement number 18 of the sequence lists.)

Formula 18

P. 238

30. New DNA which is recorded in claim article 23 in which the aforementioned DNA contains the base sequences which are recorded in the following formula 19 (arrangement number 19 of the sequence lists.)

Formula 19

P. 240

31. New DNA which is recorded in claim article 23 in which the aforementioned DNA contains the base sequences which are recorded in the following formula 20 (arrangement number 20 of the sequence lists.)

Formula 20

P. 242

32. New DNA which is recorded in claim article 23 in which the aforementioned DNA contains the base sequences which are recorded in the following formula 21 (arrangement number 21 of the sequence lists.)

Formula 21

P. 244

33. New DNA which is recorded in claim article 23 in which the aforementioned DNA contains the base sequences which are recorded in the following formula 22 (arrangement number 22 of the sequence lists.)

Formula 22

P. 245

34. New DNA which is recorded in claim article 23 in which the aforementioned DNA contains the base sequences which are recorded in the following formula 23 (arrangement number 23 of the sequence lists.)

Formula 23

P. 247

35. New DNA which is recorded in claim article 23 in which the aforementioned DNA contains the base sequences which are recorded in the following formula 24 (arrangement number 24 of the sequence lists.)

Formula 24

P. 250

36. New DNA which is recorded in claim article 23 in which the aforementioned DNA is a part of the base sequences recorded in the prior formula 16 (arrangement number 16 of the sequence lists.)

37. New DNA which is recorded in claim article 23 in which the aforementioned DNA is a part of the base sequences recorded in the prior formula 20 (arrangement

number 20 of the sequence lists.)

38. New DNA which is recorded in claim article 23 in which the aforementioned DNA is a part of the base sequences recorded in the prior formula 24 (arrangement number 24 of the sequence lists.)

39. New DNA which has the characteristic of hybridizing with DNA which contains the complementary base sequences of the base sequences recorded in any or some of prior formulas 13, 17, and 21 and coding Fas ligand (arrangement numbers 13, 17, and 21 of the sequence lists)

40. Recombinant DNA molecules which contain the new DNA recorded in either claim articles 23 or 39.

41. Transformant which has the characteristic of being transformed by the new DNA recorded in either of claim articles 23 or 39.

42. Transformant which has the characteristic of being transformed by the recombinant DNA molecules recorded in claim article 40.

43. Transformant recorded in either of claim articles 41 or 42 in which transformant transformed at least one of the hosts chosen from the following a) or c)

- a) E. coli
- b) yeast
- c) mammalian cells

P. 250-1

44. Production methods of the new polypeptides recorded in either of claim articles 1 or 22 which have the characteristic of culturing the transformant recorded in either of claim articles 41 or 43, collecting and refining new polypeptides recorded in either of claim articles 1 or 22 from the culture mixtures.

P. 251

45. Refining methods for the new polypeptides recorded in either of claim articles 1 or 22 which is a refining method for the aforementioned new polypeptides from sample which contains new polypeptides recorded in either of claim articles 1 or 22, and have the characteristic of at least performing one or more of the refining processes chosen from the following:

- (1) Affinity chromatography using Fas antigen;
- (2) Affinity chromatography using antibody which recognizes new polypeptides recorded in either of claim articles 1 or 22.

46. New antibody which has the characteristic of recognizing new polypeptides recorded in either claim articles 1 or 2.

47. New antibody recorded in claim article 46 in which the aforementioned antibody

is monoclonal antibody.

48. New antibody recorded in claim article 46 in which the aforementioned antibody is polyclonal antibody.

49. New antibodies recorded in either claim articles 46 or 48 which have the effect of suppressing apoptosis of the cells which express Fas antigen.

50. Oligonucleotides or oligonucleotide derivatives which have the characteristic of containing complementary base sequences to part of the mRNA for part of the Fas ligand gene or Fas ligand.

51. The oligonucleotides or oligonucleotide derivatives recorded in claim article 50 which control the expression of Fas ligand.

P. 251-2

52. The oligonucleotides or oligonucleotide derivatives recorded in claim articles 50 or 51 which suppress the expression of Fas ligand.

P. 252

53. Oligonucleotides or oligonucleotide derivatives recorded in either claim articles 50 or 52 in which oligonucleotides or oligonucleotide derivatives are the oligonucleotides or their derivatives which contain part of the complementary base sequences to the base sequences recorded in either of arrangement numbers 13 or 28 of the sequence lists.

54. Screening methods for the substances related to Fas ligand or the substances related to Fas antigen which have the characteristic of utilizing new polypeptides recorded in either of claim articles 1 or 22, or transformant which expresses the new polypeptides.

55. Screening methods in claim article 54 which have the characteristic of utilizing new polypeptides recorded in either of claim articles 1 or 22, or transformant and Fas antigen which expresses the new polypeptides, or cells which express Fas antigen.

56. Screening methods recorded in claim articles 54 or 55 which screen at least 1 substance chosen from the group made from substances which combine with Fas ligand, substances which control the expression of Fas ligand, substances which control the expression of Fas antigen, and substances which control the operation of Fas ligand target cells.

57. Screening methods of either of claim articles 54 or 56 which contain the process of being selected from any of the following (1) through (3).

(1) a. Culture the cells which express Fas antigen with one of polypeptides recorded in either of claim articles 1 or 22, transformants which

express the polypeptides, or groups made from the growth medium of the transformants.

b. Add the experimental substance or the sample containing the experimental substance to a.

c. Evaluate for at least one of the following: life or death of the cells which express Fas antigen; morphological changes; or biochemical changes.

P. 253

(2) a. Incubate the experimental substance or the sample containing the experimental substance with any or some of the following: new polypeptides recorded in either of claim articles 1 or 22; transformants which express the new polypeptides; or groups made from the growth medium of the transformant.

b. Culture by adding the cells which express Fas antigen to a.

c. Evaluate for at least one of the following: life or death of the cells which express Fas antigen; morphological changes; or biochemical changes.

(3) a. Incubate the experimental substance or the sample containing the experimental substance with the cells which express Fas antigen.

b. Culture by adding any or some of the following to a: new polypeptides recorded in either of prior formulas 1 or 22; transformant which expresses the new polypeptides; or groups made from the growth medium of the transformant.

c. Evaluate for at least one of the following: life or death of the cells which express Fas antigen; morphological changes; or biochemical changes.